

# Create Data Frame with Relative Weight and Gabelhouse Length Categories

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Here I need to make two data files. Both need to contain fish caught in the years 2013-2016. Both will contain the Relative Weight (Wr) of each fish and the gabel house length category each fish fits into. Then I will create two CLEAN data files one of wich will contain only fish larger than stock length and another with all fish of any length.

The data file with fish of all lengths will be used to compare the length frequency distribution between years. The data frame with only fish stock length and larger will be used to compare the proportional size densities between years and with the relative weight between years.

```
Cond <- read.csv("Data/Raw-Data/Nearshore-Biodat_2013-2017.csv") %>%
  filterD(Species == 317) %>%
  filterD(!is.na(Length)) %>%
  filterD(!is.na(Weight)) %>%
  dplyr::select(Year:Length, Age:Sex)

Cond %<>% mutate(lcat20=lencat(Length,w=20)) %>%
  mutate(logW=log10(Weight),logL=log10(Length))

Cond$fyr <- factor(Cond$Year)

Cond$fyr <- str_sub(Cond$fyr,start = 3,end = 4)
Cond %<>% dplyr::select(fyr,Year:Length,Age:logL)

#str(Cond)
headtail(Cond)
```

```
##      fyr Year Site FID Weight Length Age SexCon Sex lcat20      logW
## 1      13 2013   2  77    54   154   1     1   1   140 1.732394
## 2      13 2013   2  78    57   159   1     6   2   140 1.755875
## 3      13 2013   2  71    72   164   2     8   2   160 1.857332
## 508     17 2017  18 NA    950   385 NA     NA  NA   380 2.977724
## 509     17 2017  18 NA    900   402 NA     NA  NA   400 2.954243
## 510     17 2017  18 NA   1400   438 NA     NA  NA   420 3.146128
##              logL
## 1      2.187521
## 2      2.201397
## 3      2.214844
## 508    2.585461
## 509    2.604226
## 510    2.641474
```

```
unique(Cond$Year)
```

```
## [1] 2013 2014 2015 2016 2017
```

**Note** I found an outlier when going through '2014-2016\_condition\_largemouth-bass.Rmd' which I will now remove.

```
nrow(Cond)
(Where.r.u <- Cond[Cond$Weight==1714,]) ## row 78
(rm.outlier <- as.numeric(row.names(Cond[Cond$Weight==1714,])))

Cond <- Cond[-c(rm.outlier),] %>%
  filter(!is.na(Year))
str(Cond)
headtail(Cond)
```

Year	Number of Fish	Number of Sites	Unique
2013	113	8	~1 site (NA)
2014	143	11	2 unique (1 & 16)
2015	80	9	1 Unique (5)
2016	131	10	1 Unique (14)
2017	42	8	2 unique (3 & 9)
Total	511		6 maybe 7 Unique Sites

12 nearshore sites were sampled annually 2013 - 2017, 10 sites are sampled every year in addition to 2 sites which are sampled every five years. Length and weight data used in our analysis were obtained from 114 largemouth bass caught during 2013, 143 during 2014, 80 during 2015, and 132 during 2016. During 2013 - 2016 a total of 469 largemouth bass were obtained from 18 sites.

### Calculate Standard and Relative Weights

```
(wsLMB <- wsVal("Largemouth Bass", simplify = TRUE))
(wsLMB_min <- wsLMB[["min.TL"]])
(wsLMB_int <- wsLMB[["int"]])
(wsLMB_slp <- wsLMB[["slope"]])
```

```
Cond %<>% mutate(Ws = 10^(wsLMB_int+wsLMB_slp*logL),
  Wr=(Weight/Ws)*100)
headtail(Cond)
```

```
##      fyr Year Site FID Weight Length Age SexCon Sex lcat20      logW
## 1      13 2013   2  77    54   154   1     1   1   140 1.732394
## 2      13 2013   2  78    57   159   1     6   2   140 1.755875
## 3      13 2013   2  71    72   164   2     8   2   160 1.857332
## 507     17 2017  18  NA   950   385  NA    NA  NA   380 2.977724
## 508     17 2017  18  NA   900   402  NA    NA  NA   400 2.954243
## 509     17 2017  18  NA  1400   438  NA    NA  NA   420 3.146128
##      logL      Ws      Wr
## 1  2.187521 42.83071 126.07775
## 2  2.201397 47.55244 119.86767
## 3  2.214844 52.62401 136.81968
## 507 2.585461 859.43486 110.53775
## 508 2.604226 989.99243  90.90979
## 509 2.641474 1310.82515 106.80296
```

## Creating data file with all size fish

```
### creating size breaks for Gabelhouse Length categories for Largemouth Bass
(lmb.cuts2 <- psdVal("Largemouth Bass"))

##   substock    stock    quality preferred memorable    trophy
##         0        200        300        380        510        630

### adding gcat variable to data frame
lmb <- Cond %>%
  mutate(gcat=lencat(Length, breaks = lmb.cuts2,
                     use.names = TRUE, drop.levels = TRUE)) ### create Gabelhouse Length Categories
lmb %<>% dplyr::select(fyr:Weight, Ws, Wr, Length, lcat20, gcat, Age:Sex)
#dplyr::select(fyr:Sex, Ws, Wr, lcat20, gcat)

headtail(lmb, n=2)

##   fyr Year Site FID Weight      Ws      Wr Length lcat20    gcat
## 1   13 2013   2  77    54  42.83071 126.07775   154   140 substock
## 2   13 2013   2  78    57  47.55244 119.86767   159   140 substock
## 508  17 2017  18 NA    900 989.99243  90.90979   402   400 preferred
## 509  17 2017  18 NA   1400 1310.82515 106.80296   438   420 preferred
##   Age SexCon Sex
## 1     1     1  1
## 2     1     6  2
## 508  NA    NA  NA
## 509  NA    NA  NA

#2-9-2018#write.csv(lmb,file="Data/Clean-Data/largemouth-bass_Wr_NS.csv", row.names = FALSE)
```

## Creating Data File with Only Stock and Larger Fish

```
### adding gcat variable to data frame
Stock <- Cond %>%
  filter(Length>=lmb.cuts2["stock"]) %>%
  mutate(gcat=lencat(Length, breaks = lmb.cuts2,
                     use.names = TRUE, drop.levels = TRUE)) ### create Gabelhouse Length Categories

Stock %<>% dplyr::select(fyr:Weight, Ws, Wr, Length, lcat20, gcat, Age:Sex)

headtail(Stock, n=2)

##   fyr Year Site FID Weight      Ws      Wr Length lcat20    gcat
## 1   13 2013   2  76    126 100.7561 125.05451   200   200 stock
## 2   13 2013   2  97    154 109.2372 140.97767   205   200 stock
## 444  17 2017  18 NA    900 989.9924  90.90979   402   400 preferred
## 445  17 2017  18 NA   1400 1310.8251 106.80296   438   420 preferred
##   Age SexCon Sex
## 1     2     8  2
## 2     2     1  1
## 444  NA    NA  NA
## 445  NA    NA  NA
```

```
#2-9-2018#write.csv(Stock,file="Data/Clean-Data/largemouth-bass_Wr_Stock.csv", row.names = FALSE)
```

## Creating a Data File to Summarize Relative Weight by Year for Stock Length Individuals

```
Stock %<>% filterD(!is.na(Wr))
```

```
Summarize(Wr~fyr, data=Stock, digits = 0) ### Wr Weight by Year
```

```
##   fyr   n mean sd min  Q1 median  Q3 max
## 1  13  97 113 13  76 104   113 122 144
## 2  14 140 110 16  80  99   107 118 151
## 3  15  67 110 16  78  99   109 121 150
## 4  16 106 115 14  62 108   116 125 146
## 5  17  35 124 34  71 104   111 131 215
```

```
(Wr.fyr.gcat_Stock <- Summarize(Wr~fyr*gcat, data=Stock) %>%
  arrange(fyr,gcat))
```

```
##   fyr   gcat   n    mean      sd    min    Q1 median  Q3    max
## 1  13 preferred 16 106.85136 11.405237 92.32  98.65 104.40 114.4 133.1
## 2  13  quality 40 108.20201 11.400456 76.14 103.60 107.70 116.6 131.3
## 3  13   stock 41 120.98826 12.034098 94.13 113.20 120.80 128.4 143.8
## 4  14 preferred 18  97.67045  8.942296 83.65  90.40  98.58 103.0 115.5
## 5  14  quality 57 103.51170 11.643284 80.40  96.07 102.20 111.6 133.1
## 6  14   stock 65 118.27433 15.782376 88.74 106.70 116.10 127.5 151.3
## 7  15 preferred 15  97.09075 12.451362 78.22  86.01  96.69 105.6 119.8
## 8  15  quality 38 109.31855 12.915921 84.02 100.80 108.10 120.5 131.0
## 9  15   stock 14 124.90379 12.564609 103.80 116.60 123.90 133.9 149.8
## 10 16 preferred 10 107.72186  7.170036 94.36 104.30 107.80 111.7 118.9
## 11 16  quality 44 111.22398 14.357351 61.76 105.50 110.50 118.9 146.2
## 12 16   stock 52 120.60206 12.540774 68.71 113.90 121.60 127.1 144.9
## 13 17 preferred  9 106.38263 12.144023 86.03 106.30 107.10 110.5 128.0
## 14 17  quality 15 109.39021 17.439429 70.94  98.39 107.20 126.8 132.5
## 15 17   stock 11 157.08177 38.820674 105.00 131.20 155.60 185.8 214.8
```

```
str(Wr.fyr.gcat_Stock)
```

```
## 'data.frame':  15 obs. of  10 variables:
## $ fyr   : chr  "13" "13" "13" "14" ...
## $ gcat  : Factor w/ 3 levels "preferred","quality",...: 1 2 3 1 2 3 1 2 3 1 ...
## $ n     : num  16 40 41 18 57 65 15 38 14 10 ...
## $ mean  : num  106.9 108.2 121 97.7 103.5 ...
## $ sd    : num  11.41 11.4 12.03 8.94 11.64 ...
## $ min   : num  92.3 76.1 94.1 83.7 80.4 ...
## $ Q1    : num  98.7 103.6 113.2 90.4 96.1 ...
## $ median: num  104.4 107.7 120.8 98.6 102.2 ...
## $ Q3    : num  114 117 128 103 112 ...
## $ max   : num  133 131 144 116 133 ...
```

```
#2-9-2018#write.csv(Wr.fyr.gcat_Stock,file = "Data/Clean-Data/summary-data/relative-weight_largemouth-bass_STOCK_Summary.csv")
```

I have created a file with the relative weight of each gabelhouse length category for each year. The file name is relative-weight\_largemouth-bass\_STOCK\_Summary.csv.

## Note

The relative weight data contains only stock length individuals. This is so that I can easily compare the relative weight of fish with PSD. This is done despite the min TL being 150 mm. I may want to summarize relative weight for 150mm and greater length individuals in the future to see if young/small fish drive down or increase Wr.

## Creating a Data File to Summarize Relative Weight by Year Length $\geq$ 150mm

```
lmb.Wr <- lmb %>%  
  filter(Length >= wsLMB_min) %>%  
  filterD(Year>=2014)  
  
(lmb.Wr.gcat <- Summarize(Wr~fyr*gcat,data = lmb.Wr,digits = 0) %>%  
  arrange(fyr,gcat))
```

##	fyr	gcat	n	mean	sd	min	Q1	median	Q3	max
## 1	14	preferred	18	98	9	84	90	99	103	116
## 2	14	quality	57	104	12	80	96	102	112	133
## 3	14	stock	65	118	16	89	107	116	128	151
## 4	14	substock	3	141	18	128	131	133	147	162
## 5	15	preferred	15	97	12	78	86	97	106	120
## 6	15	quality	38	109	13	84	101	108	120	131
## 7	15	stock	14	125	13	104	117	124	134	150
## 8	15	substock	6	161	49	113	126	149	181	243
## 9	16	preferred	10	108	7	94	104	108	112	119
## 10	16	quality	44	111	14	62	106	110	119	146
## 11	16	stock	52	121	13	69	114	122	127	145
## 12	16	substock	16	128	7	117	123	127	133	145
## 13	17	preferred	9	106	12	86	106	107	110	128
## 14	17	quality	15	109	17	71	98	107	127	132
## 15	17	stock	11	157	39	105	131	156	186	215
## 16	17	substock	3	153	105	83	93	103	189	274

```
#2-9-2018#write.csv(lmb.Wr.gcat,file = "Data/Clean-Data/summary-data/relative-weight_largemouth-bass_15
```